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BLink, Conserved Domains, Links

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1: NP 416275. Reports glutamate dehydro...[gi:16129715]
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LOCUS
                                       447 aa
                                                                   BCT 27-JAN-2009
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DBSOURCE
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            Enterobacteriaceae; Escherichia.
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            Riley, M., Abe, T., Arnaud, M.B., Berlyn, M.K., Blattner, F.R.,
            Chaudhuri, R.R., Glasner, J.D., Horiuchi, T., Keseler, I.M., Kosuge, T.,
            Mori, H., Perna, N.T., Plunkett, G. III, Rudd, K.E., Serres, M.H.,
            Thomas, G.H., Thomson, N.R., Wishart, D. and Wanner, B.L.
  TITLE
            Escherichia coli K-12: a cooperatively developed annotation
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            Nucleic Acids Res. 34 (1), 1-9 (2006)
  JOURNAL
   PUBMED
            16397293
  REMARK
            Publication Status: Online-Only
REFERENCE
            2 (residues 1 to 447)
  AUTHORS
            Blattner, F.R., Plunkett, G., Bloch, C.A., Perna, N.T., Burland, V.,
            Riley, M., Collado-Vides, J., Glasner, J.D., Rode, C.K., Mayhew, G.F.,
            Gregor, J., Davis, N.W., Kirkpatrick, H.A., Goeden, M.A., Rose, D.J.,
            Mau, B. and Shao, Y.
  TITLE
            The complete genome sequence of Escherichia coli K-12
            Science 277 (5331), 1453-1474 (1997)
  JOURNAL
   PUBMED
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REFERENCE
               (sites)
  AUTHORS
            Kang, W.K., Icho, T., Isono, S., Kitakawa, M. and Isono, K.
            Characterization of the gene rimK responsible for the addition of
  TITLE
            glutamic acid residues to the C-terminus of ribosomal protein S6 in
            Escherichia coli K12
  JOURNAL.
            Mol. Gen. Genet. 217 (2-3), 281-288 (1989)
   PUBMED
            2570347
REFERENCE
                (residues 1 to 447)
  AUTHORS
            Arnaud, M., Berlyn, M.K.B., Blattner, F.R., Galperin, M.Y.,
            Glasner, J.D., Horiuchi, T., Kosuge, T., Mori, H., Perna, N.T.,
            Plunkett, G. III, Riley, M., Rudd, K.E., Serres, M.H., Thomas, G.H. and
            Wanner, B.L.
  TITLE
            Workshop on Annotation of Escherichia coli K-12
  JOURNAL
            Unpublished
            Woods Hole, Mass., on 14-18 November 2003 (sequence corrections)
  REMARK
REFERENCE
            5 (residues 1 to 447)
  AUTHORS
            Glasner, J.D., Perna, N.T., Plunkett, G. III, Anderson, B.D.,
            Bockhorst, J., Hu, J.C., Riley, M., Rudd, K.E. and Serres, M.H.
            ASAP: Escherichia coli K-12 strain MG1655 version m56
  TITLE
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NCBI Sequence Viewer v2.0 TOURNAL. Unpublished REMARK ASAP download 10 June 2004 (annotation updates) REFERENCE 6 (residues 1 to 447) AUTHORS Hayashi, K., Morooka, N., Mori, H. and Horiuchi, T. TITLE A more accurate sequence comparison between genomes of Escherichia coli K12 W3110 and MG1655 strains JOURNAL Unpublished REMARK GenBank accessions AG613214 to AG613378 (sequence corrections) REFERENCE 7 (residues 1 to 447) AUTHORS Perna, N.T. TITLE Escherichia coli K-12 MG1655 vgiK-rfaE intergenic region, genomic sequence correction JOURNAL Unpublished REMARK GenBank accession AY605712 (sequence corrections) REFERENCE 8 (residues 1 to 447) AUTHORS Rudd, K.E. A manual approach to accurate translation start site annotation; an TITLE E. coli K-12 case study JOURNAL. Unpublished REFERENCE 9 (residues 1 to 447) CONSRIM NCBI Genome Project TITLE Direct Submission JOURNAL Submitted (27-JAN-2009) National Center for Biotechnology Information, NIH, Bethesda, MD 20894, USA REFERENCE 10 (residues 1 to 447) AUTHORS Rudd, K.E. TITLE Direct Submission JOURNAL Submitted (22-DEC-2008) Department of Biochemistry and Molecular Biology, University of Miami Miller School of Medicine, 118 Gautier Bldg., Miami, FL 33136, USA REMARK Annotation update from ecogene.org as a multi-database collaboration REFERENCE 11 (residues 1 to 447) AUTHORS Rudd, K.E. TITLE Direct Submission JOURNAL Submitted (12-MAR-2008) Department of Biochemistry and Molecular Biology, University of Miami Miller School of Medicine, 118 Gautier Bldg., Miami, FL 33136, USA REMARK Annotation update from ecogene.org as a multi-database collaboration REFERENCE 12 (residues 1 to 447) AUTHORS Rudd, K.E. TITLE Direct Submission JOURNAL Submitted (26-SEP-2007) Department of Biochemistry and Molecular Biology, University of Miami Miller School of Medicine, 118 Gautier Bldg., Miami, FL 33136, USA REMARK Annotation update from ecogene.org as a multi-database collaboration REFERENCE 13 (residues 1 to 447) AUTHORS Rudd, K.E. TITLE Direct Submission JOURNAL Submitted (24-APR-2007) Department of Biochemistry and Molecular Biology, University of Miami Miller School of Medicine, 118 Gautier Bldg., Miami, FL 33136, USA REMARK Annotation update from ecogene.org as a multi-database collaboration REFERENCE 14 (residues 1 to 447) AUTHORS Plunkett, G. III. Direct Submission TITLE JOURNAL Submitted (07-FEB-2006) Laboratory of Genetics, University of Wisconsin, 425G Henry Mall, Madison, WI 53706-1580, USA REMARK Protein updates by submitter REFERENCE 15 (residues 1 to 447)

AUTHORS

Plunkett, G. III.

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NCBI Sequence Viewer v2.0

TITLE Direct Submission
JOURNAL Submitted (10-JUN-2004) Laboratory of Genetics, University of
Wisconsin, 4256 Henry Mall, Madison, WI 53706-1580, USA
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REMARK

REFERENCE 16 (residues 1 to 447)
AOTHORS Plunkett, G. III.

TITLE Direct Submission
JOURNAL Submitted (13-0CT-1998) Laboratory of Genetics, University of

Wisconsin, 425G Henry Mall, Madison, WI 53706-1580, USA REFERENCE 17 (residues 1 to 447) AUTHORS Blattner, F.R. and Plunkett, G. III.

Sequence update by submitter

TITLE Direct Submission
JOURNAL Submitted (02-SEP-1997) Laboratory of Genetics, University of

Wisconsin, 425G Henry Mall, Madison, WI 53706-1580, USA REFERENCE 18 (residues 1 to 447)

AUTHORS Blattner, F.R. and Plunkett, G. III.
TITLE Direct Submission

JOURNAL Submitted (16-JAN-1997) Laboratory of Genetics, University of Wisconsin, 425G Henry Mall, Madison, WI 53706-1580, USA

COMMENT PROVISIONAL <u>REFSEQ</u>: This record has not yet been subject to final NCBI review. The reference sequence was derived from <u>AAC74831</u>.

Method: conceptual translation.
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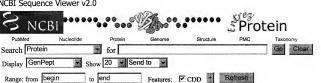
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1: AAA23868. Reports glutamate dehydro...[gi:146126]

BLink, Conserved Domains, Links

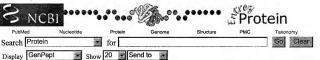


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Range: from begin





Features: CDD

to end

1: NP 288194. Reports glutamate dehydro...[gi:15802172]

BLink, Conserved Domains, Links

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ACCESSION
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            NP 288194.1 GI:15802172
VERSION
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KEYWORDS
SOURCE
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            Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
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REFERENCE
            1 (residues 1 to 447)
  AUTHORS
            Perna, N.T., Plunkett, G. III, Burland, V., Mau, B., Glasner, J.D.,
            Rose, D.J., Mayhew, G.F., Evans, P.S., Gregor, J., Kirkpatrick, H.A.,
            Posfai, G., Hackett, J., Klink, S., Boutin, A., Shao, Y., Miller, L.,
            Grotbeck, E.J., Davis, N.W., Lim, A., Dimalanta, E., Potamousis, K.,
            Apodaca, J., Anantharaman, T.S., Lin, J., Yen, G., Schwartz, D.C.,
            Welch, R.A. and Blattner, F.R.
            Genome sequence of enterohaemorrhagic Escherichia coli 0157:H7
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  JOURNAL
            Nature 409 (6819), 529-533 (2001)
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  TITLE
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  JOURNAL
            Submitted (28-SEP-2001) National Center for Biotechnology
            Information, NIH, Bethesda, MD 20894, USA
REFERENCE
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  AUTHORS
            Perna, N.T., Plunkett, G. III, Burland, V., Mau, B., Glasner, J.D.,
            Rose, D. J., Mayhew, G. F., Evans, P. S., Gregor, J., Kirkpatrick, H.A.,
            Posfai, G., Hackett, J., Klink, S., Boutin, A., Shao, Y., Miller, L.,
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            Apodaca, J., Anantharaman, T.S., Lin, J., Yen, G., Schwartz, D.C.,
            Welch, R.A. and Blattner, F.R.
  TITLE
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            Submitted (22-OCT-2000) Laboratory of Genetics, University of
  JOURNAL
            Wisconsin, 445 Henry Mall, Madison, WI 53706, USA
COMMENT
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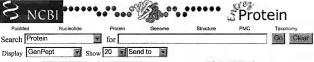
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NCBI Sequence Viewer v2.0
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Range: from begin



Features: CDD

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            NP 754056
ACCESSION
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VERSION
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REFERENCE
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  AUTHORS
            Welch, R.A., Burland, V., Plunkett, G.D. III, Redford, P., Roesch, P.,
            Rasko, D.A., Buckles, E.L., Liou, S.-R., Boutin, A., Hackett, J.,
            Stroud, D., Mayhew, G.F., Rose, D.J., Zhou, S., Schwartz, D.C.,
            Perna, N.T., Mobley, H.L.T., Donnenberg, M.S. and Blattner, F.R.
  TITLE
            Extensive mosaic structure revealed by the complete genome sequence
            of uropathogenic Escherichia coli
  JOURNAL
            Proc. Natl. Acad. Sci. U.S.A. 99 (26), 17020-17024 (2002)
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REFERENCE
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            NCBI Genome Project
  TITLE
            Direct Submission
            Submitted (10-SEP-2004) National Center for Biotechnology
  JOURNAL
            Information, NIH, Bethesda, MD 20894, USA
REFERENCE
               (residues 1 to 447)
            Welch, R.A., Burland, V., Plunkett, G.D. III, Redford, P., Roesch, P.,
  AUTHORS
            Rasko, D.A., Buckles, E.L., Liou, S.-R., Boutin, A., Hackett, J.,
            Stroud, D., Mayhew, G.F., Rose, D.J., Zhou, S., Schwartz, D.C.,
            Perna, N.T., Mobley, H.L.T., Donnenberg, M.S. and Blattner, F.R.
  TITLE
            Direct Submission
            Submitted (20-JUN-2002) Genetics Laboratory, University of
  JOURNAL
            Wisconsin - Madison, 445 Henry Mall, Madison, WI 53706, USA
COMMENT
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      241 gnvaqyaiek amefgarvit asdssgtvvd esgftkekla rlieikasrd grvadyakef
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Range: from begin

Features: CDD

Send to

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BLink, Conserved Domains, Links

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1: ZP 03049943. Reports NADP-specific glu...[gi:193068984]
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                                                                 BCT 25-JUN-2008
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REFERENCE
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  AUTHORS
            Rasko, D., Rosovitz, M., Myers, G., Seshadri, R., Cer, R., Jiang, L.,
            Ravel, J., Fricke, W.F. and Sebastian, Y.
  TITLE
            Direct Submission
  JOURNAL
            Submitted (16-JUN-2008) J. Craig Venter Institute, 9704 Medical
            Center Drive, Rockville, MD 20850, USA
REFERENCE
            2 (residues 1 to 447).
  AUTHORS
            Rasko, D.A., Rosovitz, M.J., Kaper, J.B., Myers, G.S.A., Seshadri, R.,
            Cer, R.Z., Jiang, L. and Ravel, J.
  TITLE
            Direct Submission
            Submitted (01-SEP-2005) The Institute for Genomic Research, 9712
  JOURNAL
            Medical Center Drive, Rockville, MD 20850, USA
COMMENT
            PREDICTED REFSEQ: This record has not been reviewed and the
            function is unknown. The reference sequence was derived from
            EDV88223.
            Escherichia coli E110019 was isolated from an outbreak in Finland
            (Viljanen MK et al, Lancet (1990), 336(8719): 831-4). The outbreak
            affected approximately 650 students at a school. In a fashion
            uncommon in other EPEC strains, E. coli E110019 spread from
            person-to-person to over 100 close contacts outside of the school
            setting. E. coli E110019 represents an atypical enteropathogenic E.
            coli (atypical EPEC) characterized by the lack of previously
            identified EPEC virulence factors including the EAF plasmid.
            E110019 is serotype 0111:H9.
            Method: conceptual translation.
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      121 tlpmgggkgg sdfdpkgkse gevmrfcgal mtelyrhlga dtdvpagdig vggrevqfma
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☐ 1: <u>YP_001463059</u>. Reports glutamate dehydro...[gi:157158996]

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CDS

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11

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☐ 1: <u>YP_002407302</u>. Reports glutamate dehydro...[gi:218699673]

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Comment Features Sequence LOCUS YP 002407302 447 aa linear BCT 22-DEC-2008 DEFINITION glutamate dehydrogenase, NADP-specific [Escherichia coli IAI39]. ACCESSION YP 002407302 YP 002407302.1 GI:218699673 VERSION DBSOURCE REFSEQ: accession NC 011750.1 KEYWORDS SOURCE Escherichia coli IAI39 ORGANISM Escherichia coli IAI39 Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae; Escherichia. REFERENCE (residues 1 to 447) CONSRTM NCBI Genome Project TITLE Direct Submission JOURNAL Submitted (18-DEC-2008) National Center for Biotechnology Information, NIH, Bethesda, MD 20894, USA REFERENCE (residues 1 to 447) AUTHORS Genoscope -, C.E.A. TITLE Direct Submission JOURNAL Submitted (14-DEC-2008) Genoscope - Centre National de Sequencage : BP 191 91006 EVRY cedex - FRANCE (E-mail : segref@genoscope.cns.fr - Web : www.genoscope.cns.fr) COMMENT PROVISIONAL REFSEQ: This record has not yet been subject to final NCBI review. The reference sequence was derived from CAR17428. Annotation results relative to BLAST similarities, COG assignations, enzymatic function prediction (PRIAM software), TMHMM and SignalP predictions, and synteny conservation (Syntonizer software) are available in the MaGe annotation system http://www.genoscope.cns.fr/agc/mage. Method: conceptual translation. FEATURES Location/Qualifiers 1..447 source /organism="Escherichia coli IAI39" /strain="IAI39" /db xref="taxon:585057" 1..447 Protein /product="glutamate dehydrogenase, NADP-specific" /EC_number="1.4.1.4" /function="1.3 : Glutamate family" /calculated mol wt=48424 6..447 Region /region name="PRK09414" /note="glutamate dehydrogenase; Provisional; PRK09414" /db xref="CDD:103887" 57..187 Region /region name="ELFV dehydrog N" /note="Glu/Leu/Phe/Val dehydrogenase, dimerisation domain; pfam02812"

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